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NEWS EXPRESS MAY 26 09 CURRENT WINDOWS VERSION IS V8.4, AND CURRENT DISCOVER FILE IS DATED 06 APRIL 2009.

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=> File MEDLINE, SCISEARCH, LIFESCI, BIOSIS, EMBASE, HCAPLUS, NTIS, ESBIOBASE, BIOTECHNO, WPIDS

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ENTRY SESSION
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- => S (protease or proteinase or peptidase) (4A) (variant or mutant or mutated or mutation or mutating or mutagenesis or substitution or substitute or substituted or substituting or replace or replaced or replacing or replacement or exchange or exchanged or exchanging)
- L1 14550 (PROTEASE OR PROTEINASE OR PEPTIDASE) (4A) (VARIANT OR MUTANT OR MUTATED OR MUTATION OR MUTATING OR MUTAGENESIS OR SUBSTITUTION NOR SUBSTITUTE OR SUBSTITUTED OR SUBSTITUTING OR REPLACE OR REPLACED OR REPLACING OR REPLACEMENT OR EXCHANGE OR EXCHANGED OR EXCHANGING)
- => S (protease or proteinase or peptidase) (5A) (unexpected or unpredictable (3A) cleavage or cleaved or cleaving or cleave)
- L2 15038 (PROTEASE OR PROTEINASE OR PEPTIDASE) (5A) (UNEXPECTED OR UNPRED ICTABLE (3A) CLEAVAGE OR CLEAVED OR CLEAVING OR CLEAVE)

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=> S 11 (P) 12
PROXIMITY OPERATOR LEVEL NOT CONSISTENT WITH
FIELD CODE - 'AND' OPERATOR ASSUMED 'L7 (P) L18'
PROXIMITY OPERATOR LEVEL NOT CONSISTENT WITH
FIELD CODE - 'AND' OPERATOR ASSUMED 'L9 (P) L20'
          386 L1 (P) L2
L3
=> S (protease or proteinase or peptidase) (4A) (substrate binding)
           605 (PROTEASE OR PROTEINASE OR PEPTIDASE) (4A) (SUBSTRATE BINDING)
\Rightarrow s 13 and 14
L5
            0 L3 AND L4
=> S (protease or proteinase or peptidase) (4A) (single substitution)
            8 (PROTEASE OR PROTEINASE OR PEPTIDASE) (4A) (SINGLE SUBSTITUTION)
L6
=> S (protease or proteinase or peptidase) (5A) (unexpected or unpredictable (3A)
cleavage or cleaved or cleaving or cleave)
         15038 (PROTEASE OR PROTEINASE OR PEPTIDASE) (5A) (UNEXPECTED OR UNPRED
               ICTABLE (3A) CLEAVAGE OR CLEAVED OR CLEAVING OR CLEAVE)
=> s 16 and 17
Γ8
            0 L6 AND L7
=> S (protease or proteinase or peptidase) (5A) (unexpected or unpredictable (3A)
cleavage or cleaved or cleaving or cleave)
         15038 (PROTEASE OR PROTEINASE OR PEPTIDASE) (5A) (UNEXPECTED OR UNPRED
L9
              ICTABLE (3A) CLEAVAGE OR CLEAVED OR CLEAVING OR CLEAVE)
=> S (protease or proteinase or peptidase) (3A) (variant or mutant or mutated or
mutation or mutating or mutagenesis or substitution or substitute or substituted or
substituting or replace or replaced or replacing or replacement or exchange or
exchanged or exchanging)
         11665 (PROTEASE OR PROTEINASE OR PEPTIDASE) (3A) (VARIANT OR MUTANT
              OR MUTATED OR MUTATION OR MUTATING OR MUTAGENESIS OR SUBSTITUTIO
              N OR SUBSTITUTE OR SUBSTITUTED OR SUBSTITUTING OR REPLACE OR
               REPLACED OR REPLACING OR REPLACEMENT OR EXCHANGE OR EXCHANGED
              OR EXCHANGING)
=> S (protease or proteinase or peptidase) (4A) (unexpected or unpredictable (2A)
cleavage or cleaved or cleaving or cleave)
         13404 (PROTEASE OR PROTEINASE OR PEPTIDASE) (4A) (UNEXPECTED OR UNPRED
L11
               ICTABLE (2A) CLEAVAGE OR CLEAVED OR CLEAVING OR CLEAVE)
=> s 110 and 111
L12
          315 L10 AND L11
=> duplicate
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ENTER L# LIST OR (END):112
DUPLICATE PREFERENCE IS 'MEDLINE, SCISEARCH, LIFESCI, BIOSIS, EMBASE, HCAPLUS,
ESBIOBASE, BIOTECHNO, WPIDS'
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PROCESSING COMPLETED FOR L12
           140 DUPLICATE REMOVE L12 (175 DUPLICATES REMOVED)
L13
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L13 ANSWER 1 OF 140 WPIDS COPYRIGHT 2009 THOMSON REUTERS on STN
    2009-H28271 [30] WPIDS
AN
    2009-Н49158; 2009-Н50191
CR
    Producing a stabilized, protease resistant apolipoprotein A1 (ApoA1)
TΙ
    protein variant, comprises modifying the ApoAl protein either by amino
     acid substitution or by chemical modification, and analyzing the
    proteolytic cleavage
DC
    B04; D16; S03
IN EYCKERMAN S; KAS K; LABEUR C
PΑ
    (PRON-N) PRONOTA NV
CYC 122
PIA WO 2009050275 A1 20090423 (200930) * EN 43[3]
ADT WO 2009050275 A1 WO 2008-EP64054 20081017
PRAI EP 2007-118859
                         20071019
L13 ANSWER 2 OF 140 WPIDS COPYRIGHT 2009
                                              THOMSON REUTERS on STN
     2009-F31009 [18] WPIDS
AN
CR
     2009-F36129; 2009-M04592
ΤT
    Modified polypeptide capable of lysing bacterial cell walls, useful as a
    medicament or diagnostic agent, has amino acid substitutions at protease
     cleavage sites that inhibit degradation by proteases
DC
     B04; D13; D16; D21
ΙN
    FORCHHEIM M; GRALLERT H
    (PROF-N) PROFOS AG
PA
CYC 122
                   A2 20090226 (200918)* DE 50[12]
PIA
    WO 2009024142
     WO 2009024142
                   A3 20090618 (200940) EN
     DE 102007061929 A1 20090625 (200942) DE
    WO 2009024142 A2 WO 2008-DE1378 20080819; WO 2009024142 A3 WO 2008-DE1378
ADT
     20080819; DE 102007061929 A1 DE 2007-102007061929 20071221
PRAI US 2007-957351P
                         20070822
    EP 2007-114785
                         20070822
     DE 2007-102007061929 20071221
     US 2008-32211P
                         20080228
     EP 2008-152096
                         20080228
     DE 2008-102008023448 20080514
L13 ANSWER 3 OF 140 WPIDS COPYRIGHT 2009 THOMSON REUTERS on STN
ΑN
    2009-F15981 [16]
                       WPIDS
CR
     2009-M96375; 2009-Q46329
ΤI
    New composition comprises an antigen and a heterologous hepatitis C virus
     (HCV) NS3 protease cleavage site, useful for enhancing an immune response
     to a hepatitis C antigen and for treating and preventing HCV infection
DC
    B04; C06; D16
    FRELIN L; SALLBERG M; SODERHOLM J; FELIN L
ΙN
    (TRIP-N) TRIPEP AB
PA
CYC 122
PIA WO 2009022236
                    A2 20090219 (200916)* EN 278[24]
     US 20090074803 A1 20090319 (200921)
                                          EN
     WO 2009022236 A8 20091001 (200964)
                                          EN
    WO 2009022236 A2 WO 2008-IB3047 20080815; US 20090074803 A1 Provisional US
     2007-956326P 20070816; US 20090074803 A1 Provisional US 2008-47076P
     20080422; US 20090074803 A1 US 2008-192776 20080815; WO 2009022236 A8 WO
     2008-IB3047 20080815
PRAI US 2007-956326P
                         20070816
     US 2008-47076P
                         20080422
                        20080815
     US 2008-192776
```

DUPLICATE 1

L13 ANSWER 4 OF 140 MEDLINE on STN

- 2009671757 IN-PROCESS ΜA
- PubMed ID: 19556225 DN
- Insights into the enzyme-substrate interaction in the norovirus 3C-like TΤ protease.
- ΑU Someya Yuichi; Takeda Naokazu
- Department of Virology II, National Institute of Infectious Diseases, CS 4-7-1 Gakuen, Musashi-Murayama, Tokyo 208-0011, Japan.. someya@nih.go.jp
- SO Journal of biochemistry, (2009 Oct) Vol. 146, No. 4, pp. 509-21. Electronic Publication: 2009-06-24. Journal code: 0376600. E-ISSN: 1756-2651. L-ISSN: 0021-924X.
- CY England: United Kingdom
- Journal; Article; (JOURNAL ARTICLE) (RESEARCH SUPPORT, NON-U.S. GOV'T)
- LA English
- NONMEDLINE; IN-PROCESS; NONINDEXED; Priority Journals FS
- ED Entered STN: 8 Oct 2009 Last Updated on STN: 16 Dec 2009
- L13 ANSWER 5 OF 140 HCAPLUS COPYRIGHT 2009 ACS on STN
- 2008:163673 HCAPLUS ΑN
- DN 148:231729
- Methods for engineering and synthesis of single-chain, activatable TΙ Clostridial neurotoxins comprising a functional binding domain, translocation domain, therapeutic element and exogenous protease cleavage site for use in therapy
- Steward, Lance E.; Francis, Joseph; Fernandez-Salas, Ester; Gilmore, ΙN Marcella A.; Li, Shengwen; Dolly, J. Oliver; Aoki, Kei Roger
- PΑ Allergan, Inc., USA
- SO U.S. Pat. Appl. Publ., 169pp., Cont.-in-part of U.S. Ser. No. 326,265. CODEN: USXXCO
- DT Patent
- English LA
- FAN.CNT 10

r An.	_	TENT	NO.			KIND		DATE			APPLICATION NO.				DATE				
PI	ΕP	20080032930 1700918 1700918				A1 A2 A3		20080207 20060913 20070905			US 2007-832173 EP 2006-2253					20070801 20000825			
		R:	ΑT,	BE,	CH,	DE,	DK,	ES,	FR,	GB,	GF	₹, I	Τ,	LI,	LU,	NL,	SE,	MC,	PT,
		IE, SI, LT			LT,	LV,	FI,	RO,	CY,	Y, AL									
		7132259 20060099672				В1		20061107			US	200	2000-648692				20000825		
	US							2006	0511		US	2006-326265				2	105		
		7419	-		В2		2008												
		S 20070259401 S 7422877				A1		2007			US	2006-61044			40		20061213		
						В2		2008											
	0.0	20080081355				A1		2008		US 2007-782112					20070724				
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		20080221012 20080182294				A1		20080911			US 2007-845252					20070827			
						A1		2008				2007-926812			20071029				
			0087			A1		2009				200						0800	
		US 20080311622 US 20090005313 US 20090069238				A1		2008		US 2008-182801						20080730			
						A1			20090101			2008-192419				20080815			
						A1 A1		2009				200						0800	
			20090081730					20090326				5 2008-193527				20080818			
			0030			A1		2009				200						0800	
			0030			A1		2009				200	_					0800	
		20090042270			A1		2009	-		US	200	8-1	963	81		21	0800	822	
PRAI		1999-150710P			Р		1999												
			-648			А3		2000											
			326			A2		2006											
	EΡ	2000	-964	920		A3		2000	0825										

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US 2006-610440
                       A1
                               20061213
     US 2007-782112
                         A1
                               20070724
     US 2007-829475
                         B1
                               20070727
                               20070801
     US 2007-832173
                         Α1
     US 2007-833720
                         B1
                               20070803
                         В1
     US 2007-844899
                               20070824
ASSIGNMENT HISTORY FOR US PATENT AVAILABLE IN LSUS DISPLAY FORMAT
L13 ANSWER 6 OF 140 WPIDS COPYRIGHT 2009
                                               THOMSON REUTERS on STN
     2008-020672 [82]
                       WPIDS
DNC C2008-457043 [82]
DNN N2009-048400 [82]
ΤI
    New computer system comprises directed by software correlating the
     presence of mutation in HIV-1 protease cleavage sites
     in the gag region, useful for evaluating the effectiveness of a protease
     inhibitor as an antiviral therapy against HIV
     B04; D16; S03; T01
DC
     DE MEYER S; DIERYNCK I
ΤN
     (TIBO-N) TIBOTEC PHARM LTD
PΑ
CYC 121
PIA WO 2008145606
                   A1 20081204 (200882)* EN 20[0]
                   A1 20081204 (200978)
     AU 2008257703
                                         ΕN
ADT
     WO 2008145606 A1 WO 2008-EP56356 20080523; AU 2008257703 A1 AU 2008-257703
     20080523
    AU 2008257703
                   A1 Based on WO 2008145606
PRAI EP 2007-108899
                         20070525
L13 ANSWER 7 OF 140 WPIDS COPYRIGHT 2009 THOMSON REUTERS on STN
AN
     2008-M32158 [72]
                       WPIDS
DNC C2008-376721 [72]
DNN N2008-906996 [72]
ΤI
     Identifying modified proteases with modified substrate specificity or
     other properties by contacting a collection of proteases with a protease
     trap polypeptide and identifying or selecting a protease
DC
     B04; D16; S03
IN
    MADISON E L; MADISON E
    (CATA-N) CATALYST BIOSCIENCES INC; (TORR-N) TORREY PINES INST MOLECULAR
PA
     STUDIES; (MADI-I) MADISON E L
CYC 122
PIA WO 2008045148 A2 20080417 (200872)* EN
                                              257[1]
     WO 2008045148 A3 20081016 (200872) EN
     WO 2008045148 A8 20080904 (200872) EN
     WO 2008045148 A9 20080529 (200872) EN
     TW 2008017517
                   A 20080416 (200921) ZH
    EP 2046951
                   A2 20090415 (200926) EN
    KR 2009031936
                   A 20090330 (200927) KO
                   A 20090406 (200931) NO
    NO 2008005408
     US 20090123452 A1 20090514 (200933) EN
     IN 2009CN00541 P4 20090605 (200951) EN
     AU 2007307260
                    A1 20080417 (200952)
                                          EΝ
                    A1 20080417 (200953)
     CA 2656531
                                          EΝ
                    A 20090826 (200959)
     CN 101517074
                                          ZH
                    A1 20090228 (200962)
    MX 2008016221
                                          ES
     JP 2009542218
                   W 20091203 (200979)
                                         JA 225
    WO 2008045148 A2 WO 2007-US15571 20070705; US 20090123452 A1 Provisional
     US 2006-818804P 20060705; US 20090123452 A1 Provisional US 2006-818910P
     20060705; AU 2007307260 A1 AU 2007-307260 20070705; CA 2656531 A1 CA
     2007-2656531 20070705; CN 101517074 A CN 2007-80032858 20070705; EP
     2046951 A2 EP 2007-861330 20070705; TW 2008017517 A TW 2007-124475
     20070705; US 20090123452 A1 US 2007-825627 20070705; EP 2046951 A2 PCT
     Application WO 2007-US15571 20070705; KR 2009031936 A PCT Application WO
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2007-US15571 20070705; NO 2008005408 A PCT Application WO 2007-US15571
     20070705; IN 2009CN00541 P4 PCT Application WO 2007-US15571 20070705; CA
     2656531 A1 PCT Application WO 2007-US15571 20070705; CN 101517074 A PCT
     Application WO 2007-US15571 20070705; MX 2008016221 A1 PCT Application WO
     2007-US15571 20070705; CA 2656531 A1 PCT Nat. Entry CA 2007-2656531
     20081230; MX 2008016221 A1 MX 2008-16221 20081217; NO 2008005408 A NO
     2008-5408 20081230; IN 2009CN00541 P4 IN 2009-CN541 20090129; KR
     2009031936 A KR 2009-702442 20090205; JP 2009542218 W PCT Application WO
     2007-US15571 20070705; JP 2009542218 W JP 2009-518386 20070705
FDT EP 2046951
                    A2 Based on WO 2008045148
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     WO 2008045148
                    A; AU 2007307260
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     2656531
                                                              A Based on WO
                                   Al Based on WO 2008045148
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                W Based on WO 2008045148
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PRAI US 2006-818910P
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                         20060705
     US 2007-825627
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     US 2006-818804P
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     US 2006-818910P
                         20060705
L13 ANSWER 8 OF 140 WPIDS COPYRIGHT 2009
                                              THOMSON REUTERS on STN
     2008-F49690 [36] WPIDS
     2009-E04174
    New insulin analog that comprises at least two hydrophobic amino acids
     substituted with hydrophilic amino acids, within or in close proximity to
     protease cleavage sites of parent insulin, useful for treating e.g.
     diabetes
    B04; D13; D16
    BALSCHMIDT P; HAVELUND S; HUBALEK F; LAUTRUP-LARSEN I; LUDVIGSEN S;
     NIELSEN P K; NORGAARD P; RIBEL-MADSEN U; NOERGAARD P
     (NOVO-C) NOVO NORDISK AS
    122
PIA WO 2008034881 A1 20080327 (200836)* EN 62[2]
     TW 2008029600 A 20080716 (200924) ZH
     NO 2009001563 A 20090420 (200933)
                                         ИО
     EP 2074141
                   A1 20090701 (200943) EN
     KR 2009071561 A 20090701 (200948)
     IN 2009DN01825 P1 20090529 (200951)
     AU 2007298919 A1 20080327 (200952) EN
     CN 101541830
                   A 20090923 (200964) ZH
    MX 2009002999 A1 20090430 (200970) ES
ADT WO 2008034881 A1 WO 2007-EP59990 20070920; AU 2007298919 A1 AU 2007-298919
     20070920; CN 101541830 A CN 2007-80043130 20070920; EP 2074141 A1 EP
     2007-820423 20070920; NO 2009001563 A PCT Application WO 2007-EP59990
     20070920; EP 2074141 A1 PCT Application WO 2007-EP59990 20070920; KR
     2009071561 A PCT Application WO 2007-EP59990 20070920; IN 2009DN01825 P1
     PCT Application WO 2007-EP59990 20070920; CN 101541830 A PCT Application
     WO 2007-EP59990 20070920; TW 2008029600 A TW 2007-135252 20070921; KR
     2009071561 A KR 2009-705790 20070920; IN 2009DN01825 P1 IN 2009-DN1825
     20090319; NO 2009001563 A NO 2009-1563 20090420; MX 2009002999 A1 PCT
     Application WO 2007-EP59990 20070920; MX 2009002999 A1 MX 2009-2999
     20090319
FDT EP 2074141
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                                              A; KR 2009071561
                                                                  A Based on
                    A; AU 2007298919 A1 Based on WO 2008034881
     WO 2008034881
                 A Based on WO 2008034881 A; MX 2009002999 A1 Based on WO
     101541830
     2008034881
PRAI EP 2006-121113
                         20060922
L13 ANSWER 9 OF 140 WPIDS COPYRIGHT 2009 THOMSON REUTERS on STN
     2008-B71099 [12] WPIDS
AN
     2008-B64222
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ΑN

CR

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PΑ

CYC

CR

New recombinant mammalian precursor protein comprises a protease site for ΤТ proteolytic cleavage and liberation of mature growth/differentiation factor 5 related protein, useful for preventing or treating neurodegenerative disorders DC B04; D16 PLOEGER F; POHL J; PLOGER F ΙN PA (BIOP-N) BIOPHARM GES BIOTECHNOLOGISCHEN ENTWICKL; (BIOP-N) BIOPHARM GES BIOTECHNOLOGISCHEN ENTWICKLUNGS CYC 121 PIA WO 2008009419 A1 20080124 (200812)\* EN EP 2043674 A1 20090408 (200929) EN CA 2657349 A1 20080124 (200977) EN JP 2009543566 W 20091210 (200981) JA 33 WO 2008009419 A1 WO 2007-EP6331 20070717; CA 2657349 A1 CA 2007-2657349 ADT 20070717; EP 2043674 A1 EP 2007-786127 20070717; EP 2043674 A1 PCT Application WO 2007-EP6331 20070717; CA 2657349 A1 PCT Application WO 2007-EP6331 20070717; CA 2657349 A1 PCT Nat. Entry CA 2007-2657349 20090109; JP 2009543566 W PCT Application WO 2007-EP6331 20070717; JP 2009543566 W JP 2009-519863 20070717 EP 2043674 Al Based on WO 2008009419 A; CA 2657349 Al Based on WO 2008009419 A; JP 2009543566 W Based on WO 2008009419 Α PRAI EP 2006-14928 20060718 L13 ANSWER 10 OF 140 HCAPLUS COPYRIGHT 2009 ACS on STN 2008:1272089 HCAPLUS 150:30225 DΝ An engineered protease that cleaves specifically after sulfated tyrosine ΤI ΑU Varadarajan, Navin; Georgiou, George; Iverson, Brent L. CS Departments of Chemical Engineering and Chemistry and Biochemistry, University of Texas, Austin, TX, 78712, USA Angewandte Chemie, International Edition (2008), 47(41), 7861-7863 SO CODEN: ACIEF5; ISSN: 1433-7851 Wiley-VCH Verlag GmbH & Co. KGaA PΒ DTJournal English LA OSC.G .3 THERE ARE 3 CAPLUS RECORDS THAT CITE THIS RECORD (3 CITINGS) RE.CNT 19 THERE ARE 19 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT L13 ANSWER 11 OF 140 MEDLINE on STN DUPLICATE 2 ΑN 2008615010 MEDLINE DN PubMed ID: 18710212 Automated molecular simulation based binding affinity calculator for TΙ ligand-bound HIV-1 proteases. Sadiq S Kashif; Wright David; Watson Simon J; Zasada Stefan J; Stoica ΑU Ileana; Coveney Peter V Centre for Computational Science, Department of Chemistry, University CS College London, London, WC1H OAJ, UK. Journal of chemical information and modeling, (2008 Sep) Vol. 48, No. 9, SO pp. 1909-19. Electronic Publication: 2008-08-19. Journal code: 101230060. ISSN: 1549-9596. United States CY DT Journal; Article; (JOURNAL ARTICLE) (RESEARCH SUPPORT, NON-U.S. GOV'T) (RESEARCH SUPPORT, U.S. GOV'T, NON-P.H.S.) LA English FS Priority Journals EM200811

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Entered STN: 23 Sep 2008

Last Updated on STN: 18 Nov 2008 Entered Medline: 17 Nov 2008

MEDLINE on STN L13 ANSWER 12 OF 140 DUPLICATE 3 2008676424 MEDITNE ΑN DN PubMed ID: 18674574 ΤI Sapovirus-like particles derived from polyprotein. ΑU Hansman Grant S; Oka Tomoichiro; Takeda Naokazu CS Department of Virology II, National Institute of Infectious Diseases, Japan.. q@nih.qo.jp Virus research, (2008 Nov) Vol. 137, No. 2, pp. 261-5. Electronic SO Publication: 2008-08-15. Journal code: 8410979. ISSN: 0168-1702. CY Netherlands DTJournal; Article; (JOURNAL ARTICLE) (RESEARCH SUPPORT, NON-U.S. GOV'T) LA English FS Priority Journals 200901 EΜ Entered STN: 23 Oct 2008 ED Last Updated on STN: 7 Jan 2009 Entered Medline: 6 Jan 2009 L13 ANSWER 13 OF 140 EMBASE COPYRIGHT (c) 2009 Elsevier B.V. All rights reserved on STN DUPLICATE 4 ΑN 2009033996 EMBASE ΤI Design of mutation-resistant HIV protease inhibitors with the substrate envelope hypothesis. Chellappan, S.; Reddy, G.S.K.K.; Ali, A. ΑU SO Chemtracts, (March 2008) Vol. 21, No. 3, pp. 103-104. ISSN: 1431-9268 CODEN: CHEMFW ΡВ Data Trace Publishing Company, 110 West Road, Ste. 227, Towson, Maryland, MD 21204-2316, United States. United States CY Journal; Article DTFS 004Microbiology: Bacteriology, Mycology, Parasitology and Virology 030 Clinical and Experimental Pharmacology 037 Drug Literature Index LA English SL English ΕD Entered STN: 6 Feb 2009 Last Updated on STN: 6 Feb 2009 L13 ANSWER 14 OF 140 WPIDS COPYRIGHT 2009 THOMSON REUTERS on STN ΑN 2007-830342 [77] WPIDS DNC C2007-286430 [77] Novel hepatocyte growth factor HGF precursor protein mutant composed of ТΤ HGF-alpha-chain or polypeptide region, HGF-beta-chain and peptide chain X, in pharmaceuticals for treating renal disorders, cancer, liver cirrhosis/skin ulcer B04; D16 DC ADACHI K; FUKUTA K; HAYATA D; MATSUMOTO K; NAKAMURA T ΙN PA(OSAU-C) UNIV OSAKA; (KRIN-N) KRINGLE PHARMA INC CYC 119 PIA WO 2007122975 A1 20071101 (200777)\* JA EP 2014676 A1 20090114 (200907) ΕN CA 2649800 A1 20071101 (200946) EN US 20090209463 A1 20090820 (200955) JP 2008512049 X 20090903 (200958) JA 29 ADT WO 2007122975 A1 WO 2007-JP57109 20070330; CA 2649800 A1 CA 2007-2649800 20070330; EP 2014676 A1 EP 2007-740545 20070330; EP 2014676 A1 PCT Application WO 2007-JP57109 20070330; CA 2649800 A1 PCT Application WO 2007-JP57109 20070330; US 20090209463 A1 PCT Application WO 2007-JP57109

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FDT EP 2014676
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                                               A; CA 2649800
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PRAI JP 2006-116498
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    2007-719364 [67]
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AN
DNC C2007-252308 [67]
    New coagulation factor X polypeptide with modified activation properties,
ТΤ
    useful for treating or preventing blood coagulation disorder, e.g.
    hemophilia
DC
    B04; D16
    HAUSER H; KALINA U; SCHULTE S; WEIMER T
ΤN
     (CSLB-N) CSL BEHRING GMBH; (ZLBB-N) ZLB BEHRING GMBH; (HAUS-I) HAUSER H;
PA
     (KALI-I) KALINA U; (SCHU-I) SCHULTE S; (WEIM-I) WEIMER T
CYC
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PIA WO 2007096116
                    A1 20070830 (200767)* EN
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    EP 1820508
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    KR 2008107385
                    A 20081210 (200915)
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    2008-224182 20080820; KR 2008107385 A KR 2008-720484 20080821; JP
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FDT EP 1991255
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L14 ANSWER 1 OF 3
                      MEDLINE on STN
     2003327749
                 MEDLINE
ΑN
    PubMed ID: 12858075
DΝ
ΤI
    An update on the pathogenesis and management of acquired thrombotic
    thrombocytopenic purpura.
ΑU
    Yarranton Helen; Machin Samuel J
    Haemostasis Research Unit, Department of Haematology, University College
CS
    London, London, UK.
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20070330; CA 2649800 A1 PCT Nat. Entry CA 2007-2649800 20081020; US

SO Current opinion in neurology, (2003 Jun) Vol. 16, No. 3, pp. 367-73. Ref: 48

Journal code: 9319162. ISSN: 1350-7540.

CY England: United Kingdom

DT Journal; Article; (JOURNAL ARTICLE)
General Review; (REVIEW)

LA English

FS Priority Journals

EM 200308

ED Entered STN: 15 Jul 2003 Last Updated on STN: 16 Aug 2003 Entered Medline: 15 Aug 2003

AΒ PURPOSE OF REVIEW: Thrombotic thrombocytopenic purpura, a clinical syndrome characterized by thrombocytopenia and microangiopathic haemolytic anaemia, was almost universally fatal until the introduction of plasma exchange therapy in the 1970s. Current outcomes have improved dramatically with the initiation of prompt plasma exchange, a treatment routinely used without any real understanding of why it is effective. RECENT FINDINGS: Recent advances suggest that a deficiency of a specific plasma metalloprotease, responsible for the physiological processing of von Willebrand factor multimers, plays a substantial role in the pathogenesis of congenital and acquired idiopathic thrombotic thrombocytopenic purpura. The von Willebrand factor-cleaving protease has now been identified as a new member of the ADAMTS family of metalloproteases, designated ADAMTS13. The acquired form of thrombotic thrombocytopenic purpura is associated with inhibitory autoantibodies against ADAMTS13, and the congenital chronic relapsing form is caused by mutations in the ADAMTS13 gene, resulting in a constitutional deficiency. Plasma exchange has been proved to be the most important therapy in thrombotic thrombocytopenic purpura, but clinical data for adjunctive therapies, such as corticosteroids, antiplatelet drugs and other immunosuppressive agents often used in combination with plasma exchange, are less well defined. SUMMARY: Recent advances in our understanding of the pathological mechanisms of thrombotic thrombocytopenic purpura not only provide a rationale for the previously empirical plasma exchange therapy (removal of the inhibitory antibodies and replacement of the deficient protease from the plasma infused), but may also help in developing more rational and targeted treatment strategies. This review discusses the clinical presentation, pathophysiology and current management of thrombotic thrombocytopenic purpura.

- L14 ANSWER 2 OF 3 HCAPLUS COPYRIGHT 2009 ACS on STN
- AN 2003:294313 HCAPLUS
- DN 139:50682
- TI TTP and ADAMTS13 mutation
- AU Fujimura, Yoshihiro
- CS Affiliated Hospital, Nara Prefectural Medical University, Japan
- SO Annual Review Ketsueki (2003) 153-162 CODEN: ARKNB7
- PB Chugai Igakusha
- DT Journal; General Review
- LA Japanese
- AB A review on von Willebrand factor (vWF) cleaving protease ADAMTS13 mutation in thrombotic thrombocytopenic purpura (TTP). The topics discussed are (1) unusually large vWF multimers in TTP; (2) vWF cleaving protease activity and its IgG type inhibitor; (3) TTP vs. Upshaw-Schulman syndrome; and (4) von Willebrand factor cleaving protease ADAMTS13 and its mutation in TTP.

- L14 ANSWER 3 OF 3 HCAPLUS COPYRIGHT 2009 ACS on STN
- AN 1987:82679 HCAPLUS
- DN 106:82679
- OREF 106:13549a,13552a
- TI Cleavage site mutant as a potential vaccine
- AU Homma, Morio

discussed.

- CS Sch. Med., Kobe Univ., Kobe, 650, Japan
- SO Concepts Viral Pathog. (1986), Volume 2, 388-93. Editor(s): Notkins, Abner Louis; Oldstone, Michael B. A. Publisher: Springer, New York, N. Y. CODEN: 52MXA4
- DT Conference; General Review
- LA English
- AB A review with 21 refs. Paramyxoviruses and influenza viruses become activated and replicate in multiple cycles when the envelope glycoprotein of the virus is cleaved by a host protease. In the absence of protease, the replication is limited to a single cycle. A protease activation mutant of Sendai virus was obtained, whose replication is restricted to a single cycle in the lung of mice, but which nevertheless, induces immunity. The availability

of such mutants for vaccines, their strengths and limitations are